

What is claimed is:

1. A process for identifying and enriching cell-specific target structures, in particular for the identification of cell-specific protein combination patterns on the surface of cells and for enriching such cells, wherein said process comprises the following steps:
- (a) depositing a heterogeneous cell mixture on one or plural surfaces with predefined structures, causing cells with corresponding target structures to become bound to such surface(s);
 - (b) removing any non-binding cells of said cell mixture from said surface(s);
 - (c) identifying the cell-specific target structures responsible for the binding of the cells to said surface(s);
 - (d) selecting and enriching cells with identical cell-specific target structures on said surface(s); and
 - (e) biochemically characterizing the target structures selected in procedural step (d).
2. The process as claimed in claim 1 wherein said heterogeneous cell mixture has been isolated from human or animal tissue or human or animal body fluids, or it consists of cultivated cells.
3. The process as claimed in one of the preceding claims wherein said surface is a human or animal tissue section

and/or endothelioid cells and/or protein chips and/or a cultivated piece of human or animal tissue.

4. The process as claimed in one of the preceding claims wherein the cell-specific target structures are identified in a process comprising the following steps:

- (I) automatically depositing a reagent solution Y1 that includes at least one marker molecule on said cell-specific target structure;
- (II) allowing the reagent solution Y1 to react, and automatically detecting at least one marker pattern of the target structure labeled with the reagent solution Y1;
- (III) removing said reagent solution Y1 before or after detecting the marker pattern, and repeating steps (I) and (II) with further reagent solutions Yn ($n = 2, 3, \dots, N$) each containing said at least one marker molecule and/or at least another marker molecule; and
- (IV) combining the marker patterns detected in step (II) to give a complex molecular combination pattern of the cell-specific target structure.

5. The process as claimed in one of the preceding claims wherein the selected target structures are biochemically characterized in procedural step e) by means of a molecule or molecular complex separation process, in particular a protein separation process.

6. The process as claimed in claim 5 wherein said protein separation process is a 2D gel electrophoresis.

7. The process as claimed in one of the preceding claims
5 wherein the following procedural step is performed after procedural step d):

 d1) conducting inhibition experiments regarding one or plural ingredients of the cell-specific target structures selected in procedural step (d) for detecting
10 a binding hierarchy of the ingredients.

8. The process as claimed in claim 7 wherein said ingredients are single or plural proteins of a cell-specific protein combination pattern.
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9. The process as claimed in claim 1 wherein the following procedural steps are performed instead of procedural step e):

- (f) automatically depositing a reagent
20 solution Y1 that includes at least one marker molecule on said selected and enriched cell-specific target structure;
- (g) allowing the reagent solution Y1 to react,
25 and automatically detecting at least one marker pattern of the target structure labeled with the reagent solution Y1;
- (h) removing said reagent solution Y1 before or after detecting the marker pattern, and repeating steps (f) and (g) with further
30 reagent solutions Yn (n = 2, 3, ..., N) each containing said at least one marker

molecule and/or at least another marker molecule; and

- (i) combining the marker patterns detected in step (g) to give a complex molecular combination pattern of the selected and enriched cell-specific target structure.

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